

1

SEQUENCE LISTING

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<110> SMEAL, TOD R.
     CALLOW, MARINELLA G.
     JALLAL, BAHIJA
     ZOZULYA, SERGEY
     GISHIZKY, MIKHAIL L.
<120> GEF-H1b: BIOMARKERS, COMPLEXES, ASSAYS AND THERAPEUTIC
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<141> 2003-07-02
<150> 60/460,053
<151> 2003-04-04
<150> 60/393,600
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- Gly His Phe Asn Asp Glu Ser Pro Leu Gly Leu Arg Arg Ile Leu Ser 85 90 95
- Gln Ser Thr Asp Ser Leu Asn Met Arg Asn Arg Thr Leu Ser Val Glu 100 105 110
- Ser Leu Ile Asp Glu Ala Glu Val Ile Tyr Ser Glu Leu Met Ser Asp 115 120 125
- Phe Glu Met Asp Glu Lys Asp Phe Ala Ala Asp Ser Trp Ser Leu Ala 130 135 140
- Val Asp Ser Ser Phe Leu Gln Gln His Lys Lys Glu Val Met Lys Gln 145 150 155 160
- Gln Asp Val Ile Tyr Glu Leu Ile Gln Thr Glu Leu His His Val Arg 165 170 175
- Thr Leu Lys Ile Met Thr Arg Leu Phe Arg Thr Gly Met Leu Glu Glu 180 185 190
- Leu His Leu Glu Pro Gly Val Val Gln Gly Leu Phe Pro Cys Val Asp 195 200 205
- Glu Leu Ser Asp Ile His Thr Arg Phe Leu Ser Gln Leu Leu Glu Arg 210 215 220
- Arg Arg Gln Ala Leu Cys Pro Gly Ser Thr Arg Asn Phe Val Ile His 225 230 235 240
- Arg Leu Gly Asp Leu Leu Ile Ser Gln Phe Ser Gly Pro Ser Ala Glu 245 250 255
- Gln Met Cys Lys Thr Tyr Ser Glu Phe Cys Ser Arg His Ser Lys Ala 260 265 270
- Leu Lys Leu Tyr Lys Glu Leu Tyr Ala Arg Asp Lys Arg Phe Gln Gln 275 280 285
- Phe Ile Arg Lys Val Thr Arg Pro Ala Val Leu Lys Arg His Gly Val 290 295 300
- Gln Glu Cys Ile Leu Leu Val Thr Gln Arg Ile Thr Lys Tyr Pro Leu 305 310 315 320
- Leu Ile Ser Arg Ile Leu Gln His Ser His Gly Ile Glu Glu Arg 325 330 335
- Gln Asp Leu Thr Thr Ala Leu Gly Leu Val Lys Glu Leu Leu Ser Asn 340 345 350

- Val Asp Glu Gly Ile Tyr Gln Leu Glu Lys Gly Ala Arg Leu Gln Glu 355 360 365
- Ile Tyr Asn Arg Met Asp Pro Arg Ala Gln Thr Pro Val Pro Gly Lys 370 375 380
- Gly Pro Phe Gly Arg Glu Glu Leu Leu Arg Arg Lys Leu Ile His Asp 385 390 395 400
- Gly Cys Leu Leu Trp Lys Thr Ala Thr Gly Arg Phe Lys Asp Val Leu 405 410 415
- Val Leu Leu Met Thr Asp Val Leu Val Phe Leu Gln Glu Lys Asp Gln
 420 425 430
- Lys Tyr Ile Phe Pro Thr Leu Asp Lys Pro Ser Val Val Ser Leu Gln 435 440 445
- Asn Leu Ile Val Arg Asp Ile Ala Asn Gln Glu Lys Gly Met Phe Leu 450 455 460
- Ile Ser Ala Ala Pro Pro Glu Met Tyr Glu Val His Thr Ala Ser Arg
 465 470 475 480
- Asp Asp Arg Ser Thr Trp Ile Arg Val Ile Gln Gln Ser Val Arg Thr 485 490 495
- Cys Pro Ser Arg Glu Asp Phe Pro Leu Ile Glu Thr Glu Asp Glu Ala 500 505 510
- Tyr Leu Arg Arg Ile Lys Met Glu Leu Gln Gln Lys Asp Arg Ala Leu
 515 520 525
- Val Glu Leu Leu Arg Glu Lys Val Gly Leu Phe Ala Glu Met Thr His 530 535 540
- Phe Gln Ala Glu Glu Asp Gly Gly Ser Gly Met Ala Leu Pro Thr Leu 545 550 555 560
- Pro Arg Gly Leu Phe Arg Ser Glu Ser Leu Glu Ser Pro Arg Gly Glu 565 570 575
- Arg Leu Leu Gln Asp Ala Ile Arg Glu Val Glu Gly Leu Lys Asp Leu
 580 585 590
- Leu Val Gly Pro Gly Val Glu Leu Leu Thr Pro Arg Glu Pro Ala
 595 600 605
- Leu Pro Leu Glu Pro Asp Ser Gly Gly Asn Thr Ser Pro Gly Val Thr 610 615 620
- Ala Asn Gly Glu Ala Arg Thr Phe Asn Gly Ser Ile Glu Leu Cys Arg 625 630 635 640

Ala Asp Ser Asp Ser Ser Gln Arg Asp Arg Asn Gly Asn Gln Leu Arg
645 650 655

Ser Pro Gln Glu Glu Ala Leu Gln Arg Leu Val Asn Leu Tyr Gly Leu 660 665 670

Leu His Gly Leu Gln Ala Ala Val Ala Gln Gln Asp Thr Leu Met Glu 675 680 685

Ala Arg Phe Pro Glu Gly Pro Glu Arg Arg Glu Lys Leu Cys Arg Ala 690 695 700

Asn Ser Arg Asp Gly Glu Ala Gly Arg Ala Gly Ala Ala Pro Val Ala 705 710 715 720

Pro Glu Lys Gln Ala Thr Glu Leu Ala Leu Leu Gln Arg Gln His Ala 725 730 735

Leu Leu Gln Glu Leu Arg Arg Cys Arg Arg Leu Gly Glu Glu Arg
740 745 750

Ala Thr Glu Ala Gly Ser Leu Glu Ala Arg Leu Arg Glu Ser Glu Gln
755 760 765

Ala Arg Ala Leu Leu Glu Arg Glu Ala Glu Glu Ala Arg Arg Gln Leu
770 775 780

Ala Ala Leu Gly Gln Thr Glu Pro Leu Pro Ala Glu Ala Pro Trp Ala 785 790 795 800

Arg Arg Pro Val Asp Pro Arg Arg Ser Leu Pro Ala Gly Asp Ala 805 810 815

Leu Tyr Leu Ser Phe Asn Pro Pro Gln Pro Ser Arg Gly Thr Asp Arg 820 825 830

Leu Asp Leu Pro Val Thr Thr Arg Ser Val His Arg Asn Phe Glu Asp 835 840 845

Arg Glu Arg Gln Glu Leu Gly Ser Pro Glu Glu Arg Leu Gln Asp Ser 850 855 860

Ser Asp Pro Asp Thr Gly Ser Glu Glu Glu Gly Ser Ser Arg Leu Ser 865 870 875 880

Pro Pro His Ser Pro Arg Gly Glu Thr Leu Ala Glu Thr Trp Thr Arg 885 890 895

Asp Phe Thr Arg Met Gln Asp Ile Pro Glu Glu Thr Glu Ser Arg Asp 900 905 910

Gly Glu Ala Val Ala Ser Glu Ser 915 920

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Ala Lys
<210> 5
<211> 10
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<210> 6
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<210> 7
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Phe Asn Pro Pro
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Leu Ala Lys

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<210> 10
<211> 31
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<210> 21
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Leu Asp Asn Phe
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Pro Gly Asp Pro Arg Ser Tyr Leu Asp
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Ser
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<223> any hydrophobic amino acid
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Pro Pro
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Pro Pro
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                  5
Pro Pro
<210> 31
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Leu Gln Leu Val Val Asp Pro Gly Asp Pro Arg Ser Tyr Leu Asp Asn 35 40 45

Phe

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Ser Tyr Leu Ser Ser Leu Ser Leu Ser Ser Ser Thr Tyr Pro Pro Pro 1 5 10 15

Ser Trp Gly Ser Ser Ser Asp Gln Gln Pro Ser Arg Val Ser His Glu 20 25 30

Gln Phe Arg Ala Ala Leu Gln Leu Val Val Ser Pro Gly Asp Pro Arg
35 40 45

Glu Tyr Leu Ala Asn Phe Ile Lys 50 55

<210> 34

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<223> Description of Artificial Sequence: Synthetic peptide

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Ile Ser Thr Ser Asn Leu Tyr Leu Pro Gln Asp Pro Thr Val Ala Lys
1 5 10 15

Gly Ala Leu Ala Gly Glu Asp Thr Gly Val Val Thr His Glu Gln Phe 20 25 30

Lys Ala Ala Leu Arg Met Val Val Asp Gln Gly Asp Pro Arg Leu Leu 35 40 45

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Leu Asp Ser Tyr Val Lys
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Pro Arg Arg Lys Ser Leu Val Gly Thr Pro Tyr Trp Met Ala Pro Glu
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Leu Val
<210> 43
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Trp Gly
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Asn Pro Pro
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Tyr Ser Val Glu Thr
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Thr Asn

<210> 48

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 1
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Asn Gly Ala Asn Arg Asp
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Tyr Pro Ser Asp Ser Phe Arg Gln Ser Leu Leu Gly Ser Arg Arg Gly
Arg Ser Ser Leu Ser Leu Ala Lys Ser Val Ser Thr Thr Asn
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Ser Glu Ile Lys Ser Ile Ser Glu Asn 35 40

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<212> PRT

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<223> Description of Artificial Sequence: Synthetic peptide

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Val Asp Pro Arg Arg Ser Leu Pro Ala Gly Asp Ala Leu Tyr Leu 20 25 30

Ser Phe Asn Pro Pro Gln Pro Ser Arg Gly Thr Asp 35

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<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic
 peptide

<400> 52

Cys Arg Arg Arg Ser Leu Pro Ala Gly Asp Ala Leu Tyr Leu Ser Phe

Asn Pro Pro